

## Sequence homology between human and mouse TIF

40	SEU10-27mtIF	MVLOKSMGRLAGTLLASCLLILAIQAGANALPVNTRE MVLOKSMGRLAGTLLASCLLILAIQAGANALPVNTRE
50	SEU10-28mtIF	NIEVSNEQQPXLVNTFLAKBSLADNNTDVRLLIGELLF RUDRSNEQQPXLVNTFLAKBSLADNNTDVRLLIGELLF
60	mtIF	PGVSUKDQCLMKQVLTTEVILEPQSDNQPAKQKQVVP PGVSUKDQCLMKQVLTTEVILEPQSDNQPAKQKQVVP
70	htIF	PGVSUKDQCLMKQVLTTEVILEPQSDNQPAKQKQVVP PGVSUKDQCLMKQVLTTEVILEPQSDNQPAKQKQVVP
80	mtIF	FITBLGNQESCHI9EDDNQCVVERLEKPRVKKGEGSGK FIRNBAPLSTCHI9EDDNLCARVYCKLPDKRKLGEGSGK
90	htIF	FITBLGNQESCHI9EDDNQCVVERLEKPRVKKGEGSGK FIRNBAPLSTCHI9EDDNLCARVYCKLPDKRKLGEGSGK
100	mtIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
110	htIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
120	mtIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
130	htIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
140	mtIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
150	htIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
160	mtIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
170	htIF	IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV
179		IKAIIGEGLDLFMSLRNACV IKAIIGEGLDLFMSLRNACV